

SEQUENCE LISTING

(1) GENERAL INFORMATION:

(i) APPLICANT: (other than US) THE WALTER AND ELIZA HALL INSTITUTE OF MEDICAL RESEARCH, COUNCIL FOR THE QUEENSLAND INSTITUTE FOR MEDICAL RESEARCH LIMITED, UNIVERSITY OF MELBOURNE
(US only) Perry F BARTLETT, Andrew W BOYD, Mirella DOTTORI, Lynne HARTLEY, Mary GALEA, George PAXINOS, Mark POLIZZOTTO, Trevor KILPATRICK, Mark MURPHY, Frank KONTGEN, Jason COONAN, Ursula GREFERATH

(ii) TITLE OF INVENTION: A METHOD OF TREATMENT

(iii) NUMBER OF SEQUENCES: 6

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: DAVIES COLLISON CAVE
(B) STREET: 1 LITTLE COLLINS STREET
(C) CITY: MELBOURNE
(D) STATE: VICTORIA
(E) COUNTRY: AUSTRALIA
(F) ZIP: 3000

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Floppy disk
(B) COMPUTER: IBM PC compatible
(C) OPERATING SYSTEM: PC-DOS/MS-DOS
(D) SOFTWARE: PatentIn Release #1.0, Version #1.25

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: INTERNATIONAL (PCT)
(B) FILING DATE: 27-OCT-1999
(C) CLASSIFICATION:

(vi) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: AUSTRALIAN PROVISIONAL
(B) FILING DATE: 27-OCT-1998
(C) CLASSIFICATION:

(viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: HUGHES, DR E JOHN L
(C) REFERENCE/DOCKET NUMBER: EJH/AF

(ix) TELECOMMUNICATION INFORMATION:

- (A) TELEPHONE: +61 3 9254 2777
- (B) TELEFAX: +61 3 9254 2770
- (C) TELEX: AA 31787

(2) INFORMATION FOR SEQ ID NO:1:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 27 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Oligonucleotides

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

CGTGCTACTT CCATTTGTCA CGTCCTG

27

(2) INFORMATION FOR SEQ ID NO:2:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 21 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Oligonucleotides

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

TGCCGTGATA GCAAATTTGA G

21

(2) INFORMATION FOR SEQ ID NO:3:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 22 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Oligonucleotides

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

AGGAAGTGAG CATTATGGAT GA

22

(2) INFORMATION FOR SEQ ID NO:4:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Oligonucleotides

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

TGCTCCTCGT GCCCAGCGTT

20

(2) INFORMATION FOR SEQ ID NO:5:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 27 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: Oligonucleotides
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

TTAGAATTCC CCGAGGAGGA GCTGTAC

27

(2) INFORMATION FOR SEQ ID NO:6:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 27 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: Oligonucleotides
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

CTAGAATTCT GCAGTCCCCAC CACCCCG

27